

SCORE Search Results Details for Application 08765588 and Search Result us-08-765-588a- 16.rni.

[Score Home Page](#)

[Retrieve Application List](#)

[SCORE System Overview](#)

[SCORE FAQ](#)

[Comments / Suggestions](#)

This page gives you Search Results detail for the Application 08765588 and Search Result us-08-765-588a-16.rni.

[start](#)

[Go Back to previous page](#)

GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.

OM nucleic - nucleic search, using sw model

Run on: March 11, 2005, 01:50:10 ; Search time 203 Seconds
(without alignments)
10011.112 Million cell updates/sec

Title: US-08-765-588A-16

Perfect score: 1242

Sequence: 1 gcacgagctcaggccgtcgc.....aaaaaaaaaaaaaaaaaa 1242

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 1202784 seqs, 818138359 residues

Total number of hits satisfying chosen parameters: 2405568

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_NA:
1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:
2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:
3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:
4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:
5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:
6: /cgn2_6/ptodata/1/ina/backfiles1.seq:
*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

%

Result No.	Score	Query Match	Length	DB	ID	Description
------------	-------	-------------	--------	----	----	-------------

1	686.4	55.3	886	1	US-08-469-427A-1	Sequence 1, Appli
2	686.4	55.3	886	2	US-08-609-443B-1	Sequence 1, Appli
3	686.4	55.3	886	2	US-08-569-063C-1	Sequence 1, Appli
4	686.4	55.3	886	3	US-08-851-896-1	Sequence 1, Appli
5	609.6	49.1	624	2	US-08-609-443B-12	Sequence 12, Appli
6	609.6	49.1	624	2	US-08-569-063C-12	Sequence 12, Appli
7	609.6	49.1	624	3	US-08-851-896-12	Sequence 12, Appli
8	547.6	44.1	1099	4	US-09-949-016-1545	Sequence 1545, Appli
9	515.2	41.5	624	2	US-08-609-443B-14	Sequence 14, Appli
10	515.2	41.5	624	2	US-08-569-063C-14	Sequence 14, Appli
11	515.2	41.5	624	3	US-08-851-896-14	Sequence 14, Appli
12	441.2	35.5	565	1	US-08-469-427A-4	Sequence 4, Appli
13	441.2	35.5	565	2	US-08-609-443B-4	Sequence 4, Appli
14	441.2	35.5	565	2	US-08-569-063C-4	Sequence 4, Appli
15	441.2	35.5	565	3	US-08-851-896-4	Sequence 4, Appli
16	412.6	33.2	591	1	US-08-469-427A-6	Sequence 6, Appli
17	412.6	33.2	591	2	US-08-609-443B-6	Sequence 6, Appli
18	412.6	33.2	591	2	US-08-569-063C-6	Sequence 6, Appli
19	412.6	33.2	591	3	US-08-851-896-6	Sequence 6, Appli
20	382.2	30.8	405	1	US-08-469-427A-8	Sequence 8, Appli
21	382.2	30.8	405	2	US-08-609-443B-8	Sequence 8, Appli
22	382.2	30.8	405	2	US-08-569-063C-8	Sequence 8, Appli
23	382.2	30.8	405	3	US-08-851-896-8	Sequence 8, Appli
24	355.6	28.6	570	1	US-08-469-427A-10	Sequence 10, Appli
25	355.6	28.6	570	2	US-08-609-443B-10	Sequence 10, Appli
26	355.6	28.6	570	2	US-08-569-063C-10	Sequence 10, Appli
27	355.6	28.6	570	3	US-08-851-896-10	Sequence 10, Appli
28	207.4	16.7	7386	4	US-09-949-016-13287	Sequence 13287, Appli
29	92.2	7.4	3583	4	US-09-976-594-921	Sequence 921, Appli
c 30	92.2	7.4	5163	4	US-09-919-039-166	Sequence 166, Appli
31	91.6	7.4	426	4	US-09-884-050-1	Sequence 1, Appli
32	91.6	7.4	444	3	US-09-392-932-6	Sequence 6, Appli
33	91.6	7.4	444	4	US-09-574-708A-1	Sequence 1, Appli
34	91.6	7.4	444	4	US-09-392-931-1	Sequence 1, Appli
35	91.6	7.4	456	5	PCT-US95-10973A-88	Sequence 88, Appli
36	91.6	7.4	467	5	PCT-US95-10973A-86	Sequence 86, Appli
37	91.6	7.4	473	3	US-08-718-904-1	Sequence 1, Appli
38	91.6	7.4	473	4	US-09-449-249-1	Sequence 1, Appli
39	91.6	7.4	473	5	PCT-US95-10973A-25	Sequence 25, Appli
40	91.6	7.4	495	4	US-09-244-583-25	Sequence 25, Appli
41	91.6	7.4	495	4	US-09-037-983C-14	Sequence 14, Appli
42	91.6	7.4	498	6	5194596-20	Patent No. 5194596
43	91.6	7.4	498	6	5194596-20	Patent No. 5194596
44	91.6	7.4	516	3	US-08-784-551C-1	Sequence 1, Appli
45	91.6	7.4	516	3	US-09-392-932-7	Sequence 7, Appli

ALIGNMENTS

RESULT 1

US-08-469-427A-1
; Sequence 1, Application US/08469427A
; Patent No. 5607918
; GENERAL INFORMATION:
; APPLICANT: Eriksson, Ulf
; APPLICANT: Olofsson, Birgitta
; APPLICANT: Alitalo, Kari
; APPLICANT: Pajusola, Katri
; TITLE OF INVENTION: VASCULAR ENDOTHELIAL GROWTH FACTOR-B AND
; TITLE OF INVENTION: DNA CODING THEREFOR

SCORE Search Results Details for Application 087 and Search Result us-08-765-588a-16.rnpb

[Score Home Page](#) [Retrieve Application List](#) [SCORE System Overview](#) [SCORE FAQ](#) [Comments / Sugg](#)

This page gives you Search Results detail for the Application 08765588 and Search Result us-08-765-588a-16.rnpb.

[start](#)

[Go Back to](#)

GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.

OM nucleic - nucleic search, using sw model

Run on: March 11, 2005, 04:17:05 ; Search time 724 Seconds
(without alignments)
10204.516 Million cell updates/sec

Title: US-08-765-588A-16

Perfect score: 1242

Sequence: 1 gcacgagctcaggccgtcgc.....aaaaaaaaaaaaaaaaaaaa 1242

Scoring table: IDENTITY_NUC

Gapop 10.0 , Gapext 1.0

Searched: 5537552 seqs, 2974263231 residues

Total number of hits satisfying chosen parameters: 11075104

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Published_Applications_NA:*

1: /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*

2: /cgn2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*

3: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*

4: /cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*

5: /cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*

6: /cgn2_6/ptodata/1/pubpna/PCTUS_PUBCOMB.seq:*

7: /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*

8: /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*

9: /cgn2_6/ptodata/1/pubpna/US09A_PUBCOMB.seq:*

10: /cgn2_6/ptodata/1/pubpna/US09B_PUBCOMB.seq:*

11: /cgn2_6/ptodata/1/pubpna/US09C_PUBCOMB.seq:*

12: /cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*

13: /cgn2_6/ptodata/1/pubpna/US10A_PUBCOMB.seq:*

14: /cgn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*

15: /cgn2_6/ptodata/1/pubpna/US10C_PUBCOMB.seq:*

16: /cgn2_6/ptodata/1/pubpna/US10D_PUBCOMB.seq:*

17: /cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq:*

18: /cgn2_6/ptodata/1/pubpna/US10F_PUBCOMB.seq:*

19: /cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*

```

20: /cgn2_6/ptodata/1/pubpna/US11_NEW_PUB.seq:*
21: /cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
22: /cgn2_6/ptodata/1/pubpna/US60_PUBCOMB.seq:*

```

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB	ID	Description
1	1242	100.0	1242	9	US-09-349-954A-16	Sequence 16, Appl
2	1242	100.0	1242	9	US-09-907-007-16	Sequence 16, Appl
3	1242	100.0	1242	17	US-10-673-708-16	Sequence 16, Appl
4	686.4	55.3	886	10	US-09-961-756-1	Sequence 1, Appl
5	623.4	50.2	1172	19	US-10-278-698-68	Sequence 68, Appl
6	623.4	50.2	1172	19	US-10-278-698-582	Sequence 582, App
7	623.4	50.2	1181	16	US-10-136-819-10	Sequence 10, Appl
8	609.6	49.1	624	10	US-09-961-756-12	Sequence 12, Appl
9	595.2	47.9	1094	9	US-09-349-954A-3	Sequence 3, Appl
10	595.2	47.9	1094	9	US-09-907-007-3	Sequence 3, Appl
11	595.2	47.9	1094	17	US-10-673-708-3	Sequence 3, Appl
12	595.2	47.9	1094	18	US-10-220-324A-1	Sequence 1, Appl
13	571.8	46.0	755	15	US-10-262-538-21	Sequence 21, Appl
14	571.8	46.0	755	15	US-10-007-926A-466	Sequence 466, App
15	571.8	46.0	755	17	US-10-174-128-4	Sequence 4, Appl
16	571.8	46.0	755	18	US-10-669-176-21	Sequence 21, Appl
17	515.2	41.5	624	9	US-09-912-436-3	Sequence 3, Appl
18	515.2	41.5	624	10	US-09-961-756-14	Sequence 14, Appl
19	515.2	41.5	624	18	US-10-384-339C-122	Sequence 122, App
20	515.2	41.5	624	18	US-10-772-927A-3	Sequence 3, Appl
21	513.4	41.3	5695	9	US-09-912-436-10	Sequence 10, Appl
22	512.6	41.3	663	9	US-09-244-694-19	Sequence 19, Appl
23	507	40.8	5695	9	US-09-912-436-9	Sequence 9, Appl
24	503.6	40.5	666	9	US-09-244-694-1	Sequence 1, Appl
25	456.8	36.8	491	9	US-09-244-694-140	Sequence 140, App
26	441.2	35.5	565	10	US-09-961-756-4	Sequence 4, Appl
27	432.8	34.8	910	9	US-09-349-954A-9	Sequence 9, Appl
28	432.8	34.8	910	9	US-09-907-007-9	Sequence 9, Appl
29	432.8	34.8	910	17	US-10-673-708-9	Sequence 9, Appl
30	412.6	33.2	591	10	US-09-961-756-6	Sequence 6, Appl
31	404.4	32.6	1154	17	US-10-264-049-846	Sequence 846, App
32	397	32.0	412	9	US-09-244-694-83	Sequence 83, Appl
33	397	32.0	415	9	US-09-244-694-84	Sequence 84, Appl
34	392.8	31.6	993	9	US-09-349-954A-5	Sequence 5, Appl
35	392.8	31.6	993	9	US-09-907-007-5	Sequence 5, Appl
36	392.8	31.6	993	17	US-10-673-708-5	Sequence 5, Appl
37	392.8	31.6	993	18	US-10-220-324A-3	Sequence 3, Appl
38	382.2	30.8	405	10	US-09-961-756-8	Sequence 8, Appl
39	375.8	30.3	410	9	US-09-244-694-141	Sequence 141, App
40	370	29.8	381	9	US-09-244-694-36	Sequence 36, Appl
41	355.6	28.6	570	9	US-09-795-006A-116	Sequence 116, App
42	355.6	28.6	570	10	US-09-961-756-10	Sequence 10, Appl
43	355.6	28.6	570	18	US-10-772-927A-1	Sequence 1, Appl
44	355.2	28.6	567	9	US-09-912-436-1	Sequence 1, Appl
45	353.2	28.4	5614	9	US-09-912-436-7	Sequence 7, Appl

ALIGNMENTS

SCORE Search Results Details for Application 08765588 and Search Result us-08-765-588a- 16.rnpm.

[Score Home Page](#)

[Retrieve Application List](#)

[SCORE System Overview](#)

[SCORE FAQ](#)

[Comments / Suggestions](#)

This page gives you Search Results detail for the Application 08765588 and Search Result us-08-765-588a-16.rnpm.

[start](#)

[Go Back to previous page](#)

GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.

OM nucleic - nucleic search, using sw model

Run on: March 11, 2005, 03:12:43 ; Search time 4480 Seconds
(without alignments)
11317.460 Million cell updates/sec

Title: US-08-765-588A-16

Perfect score: 1242

Sequence: 1 gcacgagctcaggccgtcgc.....aaaaaaaaaaaaaaaaaa 1242

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 45554873 seqs, 20411521753 residues

Total number of hits satisfying chosen parameters: 91109746

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Pending_Patents_NA_Main:*

1: /cgn2_6/ptodata/1/pna/PCTUS1_COMB.seq:*

2: /cgn2_6/ptodata/1/pna/PCTUS2_COMB.seq:*

3: /cgn2_6/ptodata/1/pna/PCTUS_COMB.seq:*

4: /cgn2_6/ptodata/1/pna/US06_COMB.seq:*

5: /cgn2_6/ptodata/1/pna/US07_COMB.seq:*

6: /cgn2_6/ptodata/1/pna/US080_COMB.seq:*

7: /cgn2_6/ptodata/1/pna/US081_COMB.seq:*

8: /cgn2_6/ptodata/1/pna/US082_COMB.seq:*

9: /cgn2_6/ptodata/1/pna/US083_COMB.seq:*

10: /cgn2_6/ptodata/1/pna/US084_COMB.seq:*

11: /cgn2_6/ptodata/1/pna/US085_COMB.seq:*

12: /cgn2_6/ptodata/1/pna/US086_COMB.seq:*

13: /cgn2_6/ptodata/1/pna/US087_COMB.seq:*

14: /cgn2_6/ptodata/1/pna/US088_COMB.seq:*

15: /cgn2_6/ptodata/1/pna/US089_COMB.seq:*

16: /cgn2_6/ptodata/1/pna/US090_COMB.seq:*

17: /cgn2_6/ptodata/1/pna/US091_COMB.seq:*

18: /cgn2_6/ptodata/1/pna/US092A_COMB.seq:*

19: /cgn2_6/ptodata/1/pna/US092B_COMB.seq:*

20: /cgn2_6/ptodata/1/pna/US093A_COMB.seq:*

21: /cgn2_6/ptodata/1/pna/US093B_COMB.seq:*

22: /cgn2_6/ptodata/1/pna/US094_COMB.seq:*

23: /cgn2_6/ptodata/1/pna/US095A_COMB.seq:*

24: /cgn2_6/ptodata/1/pna/US095B_COMB.seq:*

25: /cgn2_6/ptodata/1/pna/US095C_COMB.seq:*

26: /cgn2_6/ptodata/1/pna/US095D_COMB.seq:*

27: /cgn2_6/ptodata/1/pna/US096A_COMB.seq:*

28: /cgn2_6/ptodata/1/pna/US096B_COMB.seq:*

29: /cgn2_6/ptodata/1/pna/US096C_COMB.seq:*

30: /cgn2_6/ptodata/1/pna/US096D_COMB.seq:*

31: /cgn2_6/ptodata/1/pna/US096E_COMB.seq:*

32: /cgn2_6/ptodata/1/pna/US097A_COMB.seq:*

33: /cgn2_6/ptodata/1/pna/US097B_COMB.seq:*

34: /cgn2_6/ptodata/1/pna/US097C_COMB.seq:*

35: /cgn2_6/ptodata/1/pna/US098A_COMB.seq:*

36: /cgn2_6/ptodata/1/pna/US098B_COMB.seq:*

37: /cgn2_6/ptodata/1/pna/US098C_COMB.seq:*

38: /cgn2_6/ptodata/1/pna/US098D_COMB.seq:*

39: /cgn2_6/ptodata/1/pna/US099A_COMB.seq:*

40: /cgn2_6/ptodata/1/pna/US099B_COMB.seq:*

41: /cgn2_6/ptodata/1/pna/US099C_COMB.seq:*

42: /cgn2_6/ptodata/1/pna/US099D_COMB.seq:*

43: /cgn2_6/ptodata/1/pna/US099E_COMB.seq:*

44: /cgn2_6/ptodata/1/pna/US099F_COMB.seq:*

45: /cgn2_6/ptodata/1/pna/US099G_COMB.seq:*

46: /cgn2_6/ptodata/1/pna/US100A_COMB.seq:*

47: /cgn2_6/ptodata/1/pna/US100B_COMB.seq:*

48: /cgn2_6/ptodata/1/pna/US101A_COMB.seq:*

49: /cgn2_6/ptodata/1/pna/US101B_COMB.seq:*

50: /cgn2_6/ptodata/1/pna/US102A_COMB.seq:*

51: /cgn2_6/ptodata/1/pna/US102B_COMB.seq:*

52: /cgn2_6/ptodata/1/pna/US103A_COMB.seq:*

53: /cgn2_6/ptodata/1/pna/US103B_COMB.seq:*

54: /cgn2_6/ptodata/1/pna/US104A_COMB.seq:*

55: /cgn2_6/ptodata/1/pna/US104B_COMB.seq:*

56: /cgn2_6/ptodata/1/pna/US105A_COMB.seq:*

57: /cgn2_6/ptodata/1/pna/US105B_COMB.seq:*

58: /cgn2_6/ptodata/1/pna/US106A_COMB.seq:*

59: /cgn2_6/ptodata/1/pna/US107A_COMB.seq:*

60: /cgn2_6/ptodata/1/pna/US107B_COMB.seq:*

61: /cgn2_6/ptodata/1/pna/US107C_COMB.seq:*

62: /cgn2_6/ptodata/1/pna/US107D_COMB.seq:*

63: /cgn2_6/ptodata/1/pna/US108A_COMB.seq:*

64: /cgn2_6/ptodata/1/pna/US108B_COMB.seq:*

65: /cgn2_6/ptodata/1/pna/US109A_COMB.seq:*

66: /cgn2_6/ptodata/1/pna/US109B_COMB.seq:*

67: /cgn2_6/ptodata/1/pna/US109C_COMB.seq:*

68: /cgn2_6/ptodata/1/pna/US110_COMB.seq:*

69: /cgn2_6/ptodata/1/pna/US6000_COMB.seq:*

70: /cgn2_6/ptodata/1/pna/US6001_COMB.seq:*

71: /cgn2_6/ptodata/1/pna/US6002_COMB.seq:*

72: /cgn2_6/ptodata/1/pna/US6003_COMB.seq:*

73: /cgn2_6/ptodata/1/pna/US6004_COMB.seq:*

74: /cgn2_6/ptodata/1/pna/US6005_COMB.seq:*

75: /cgn2_6/ptodata/1/pna/US6006_COMB.seq:*

76: /cgn2_6/ptodata/1/pna/US6007_COMB.seq:*

77: /cgn2_6/ptodata/1/pna/US6008_COMB.seq:*

SCORE Search Results Details for Application 08765588 and Search Result us-08-765-588a- 16.rnnpn.

[Score Home Page](#) [Retrieve Application List](#) [SCORE System Overview](#) [SCORE FAQ](#) [Comments / Suggestions](#)

This page gives you Search Results detail for the Application 08765588 and Search Result us-08-765-588a-16.rnnpn.

[start](#)

[Go Back to previous page](#)

GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.

OM nucleic - nucleic search, using sw model

Run on: March 11, 2005, 04:13:05 ; Search time 1462 Seconds
(without alignments)
624.206 Million cell updates/sec

Title: US-08-765-588A-16

Perfect score: 1242

Sequence: 1 gcacgagctcaggccgtcgc.....aaaaaaaaaaaaaaaaaaaa 1242

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 5522993 seqs, 367386938 residues

Total number of hits satisfying chosen parameters: 11045986

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Pending_Patents_NA_New:*

1: /cgn2_6/ptodata/1/pna/PCT_NEW_COMB.seq:*

2: /cgn2_6/ptodata/1/pna/US06_NEW_COMB.seq:*

3: /cgn2_6/ptodata/1/pna/US07_NEW_COMB.seq:*

4: /cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:*

5: /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*

6: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:*

7: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq2:*

8: /cgn2_6/ptodata/1/pna/US11_NEW_COMB.seq:*

9: /cgn2_6/ptodata/1/pna/US60_NEW_COMB.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

%

Day : Thursday
Date: 6/15/2006

Time: 10:02:02

PALM INTRANET**Inventor Information for 08/765588**

Inventor Name	City	State/Country
HAYWARD, NICHOLAS KIM	PADDINGTON	AUSTRALIA
WEBER, GUNTHER	STOCKHOLM	SWEDEN
GRIMMOND, SEAN	TARINGA	AUSTRALIA
NORDENSKJOLD, MAGNUS	STOCKHOLM	SWEDEN
LARSSON, CATHARINA	STOCKHOLM	SWEDEN

[Appn Info](#)[Contents](#)[Petition Info](#)[Atty/Agent Info](#)[Continuity Data](#)[Foreign Data](#)

Search Another: Application# or Patent#
 PCT / / or PG PUBS #
 Attorney Docket #
 Bar Code #

To go back use Back button on your browser toolbar.

Back to [PALM](#) | [ASSIGNMENT](#) | [OASIS](#) | Home page

Day : Thursday
Date: 6/15/2006

Time: 10:03:13



Inventor Information for 08/569063

Inventor Name	City	State/Country
ERIKSSON, ULF	BALSTA	SWEDEN
OLOFSSON, BIRGITTA	SUNDBYBERG	SWEDEN
ALITALO, KARI	HELSINKI	FINLAND
PAJUSOLA, KATRI	HELSINKI	FINLAND

Search Another: Application# or Patent#
 PCT / / or PG PUBS #
 Attorney Docket #
 Bar Code #

To go back use Back button on your browser toolbar.

Back to [PALM](#) | [ASSIGNMENT](#) | [OASIS](#) | [Home page](#)